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Telephone Number



United States Patent [19]

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[54]	METHODS FOR SORTING
	POLYNUCLEOTIDES USING
	OLIGONUCLEOTIDE TAGS

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[73] Assignee: Spectragen, Inc., Hayward, Calif.

[21] Appl. No.: 358,810

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ted U.S. Application Data

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[51]	Int. Cl.6			
[52]	U.S. Cl	435/6; 536/25.4; 435/172.3		
[58]	Field of Searc	th		
[56]	1	References Cited		
	U.S. P	ATENT DOCUMENTS		
_	5,302,509 4/199 5,482,836 1/199			

FOREIGN PATENT DOCUMENTS

2036946	10/1991	Canada .
0303459A3	2/1989	European Pal. Off
90107066.4	10/1990	European Pat. Off
WO93/17126	9/1933	WIPO.
WO90/03382	4/1990	WIPO.
WO92/00091	1/1992	WIPO.
WO92/10587	6/1992	WIPO.
WO92/10588	6/1992	WIPO.
WO93/06121	4/1993	WIPO.
WO93/21203	10/1993	WIPO.
WO93/22680	11/1993	WIPO.
WO93/22684	11/1993	WIPO.
WO94/08051	4/1994	WIPO.
WO95/20053	7/1995	WIPO.

OTHER PUBLICATIONS

Kuijper et al, "Functional cloning vectors for use in directional cDNA cloning using cohesive ends produced with T4

DNA polymerase," Gene, 112: 147-155 (1992).

Asianidis et al, "Ligation-independent cloning of PCR products (LIC-PCR)," Nucleic Acids Research, 18: 6069-6074 (1990).

Wetmur, "DNA probes: applications of the principles of nucleic and hybridization," Critical Reviews in Biochemis-

try and Molecular Biology, 26: 227-259 (1991). Egholm et al, "PNA hybridizes to complementary oligonucleotides obeying the Watson-Crick hydrogen-bonding rules," Nature, 365: 566-568 (1993).

Oryaznov et al, "Modulation of oligonucleotide duplex and

triples stability via hydrophobic interactions," Nucleic Acids Research, 21: 5909-5915 (1993).

Crick et al, "Codes without commas," Proc. Natl. Acad.

Sci., 43: 416-421 (1957).

Ohlmeyer et al, "Complex synthetic chemical libraries indexed with molecular tags," Proc. Natl. Acad. Sci., 90: 10922-10926 (1993).

Brenner and Lerner, "Encoded combinatorial chemistry," Proc. Natl. Acad. Sci., 89: 5381-5383 (1992).

Maskos and Southern, "Oligonucleotide hybridizations on glass supports: a novel linker for oligonucleotide synthesis and hybridization properties of oligonucleotides synthesized in situ," Nucleic Acids Research, 20: 1679-1684 (1992). Matthews and Kricka, "Analytical strategies for the use of DNA probes," Anal. Biochem. 169: 1-25 (1988).

Broude et al, "Enhanced DNA sequencing by hybridization," Proc. Natl. Acad. Sci., 91; 3072-3076 (1994).

Nielsen et al, "Synthetic methods for the implementation of encoded combinatorial chemistry," J. Am. Chem. Soc. 115: 9812-9813 (1993).

Needels et al., "Generation and screening of an oligonucle-otide-encoded synthetic peptide library," Proc. Natl. Acad. Sci., 90: 10700-10704 (1993): Chetverin et al. "Oligonucleotide arrays: New concepts and

possibilities," Biotechnology, 12: 1093-1099 (1994).

Yang and Youvan, "A prospectus for multispectral-multiplex DNA sequencing," Biotechnology, 7: 576-580 (1989). Church et al, "Multiplex DNA Sequencing," Science, 240: 185-188 (1988).

Beck et al, "A Strategy for the amplification, purification, and selection of M13 templates for large-scale DNA sequencing," Analytical Biochemistry, 212: 498-505 (1993).

Ji and Smith, "Rapid purification of double-stranded DNA by triple-helix-mediated affinity capture," Anal.Chem., 65: 1323-1328 (1993).

Brown et al., "A new base-stable linker for solid-phase oligonucleotide synthesis," J. Chem. Soc. Commun. 1989:

Oliphant et al, "Cloning of random-sequence oligodeoxy-

uucleotides," Gene, 44: 177-183 (1986). Hunkapiller et al, "Large-scale and automated DNA sequence determination." Science, 254: 59-67 (1991).

Coche et al., "Reducing bias in cDNA sequence representation by molecular selection," Nucleic Acids Research, 22: 4545-4546 (1994).

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ABSTRACT

The invention provides a method of tracking, identifying, and/or sorting classes or subpopulations of molecules by the use of oligonucleotide tags. Oligonucleotide tags of the invention each consist of a plurality of subunits 3 to 6 nucleotides in length selected from a minimally crosshybridizing set. A subunit of a minimally cross-hybridizing set forms a duplex or triplex having two or more mismatches with the complement of any other subunit of the same set. The number of oligonucleotide tags available in a particular embodiment depends on the number of subunits per tag and on the length of the subunit. An important aspect of the invention is the use of the oligomucleotide tags for sorting polymucleotides by specifically hybridizing tags attached to the polynucleotides to their complements on solid phase supports. This embodiment provides a readily automated system for manipulating and sorting polynucleotides, particularly useful in large-scale parallel operations, such as large-scale DNA sequencing, mRNA fingerprinting, and the like, wherein many target polynucleotides or many segments of a single target polynucleotide are sequenced simultancously.

31 Claims, 6 Drawing Sheets